REV-01 BMB/01/04

B.Sc. MICROBIOLOGY FIFTH SEMESTER (SPECIAL REPEAT) BIOINFORMATICS BMB-503

[USE OMR SHEET FOR OBJECTIVE PART]

Duration: 3 hrs.

Time: 30 mins.

Objective)

Full Marks: 70

2023/08

Marks: 20

Choose the correct answer from the following:				
1.	Which of the following database is of short a. PROSITE c. PDB	b.	uence pattern and profile of protein? iProclass Pfam	
2.	Which of the following measures is not affe a. Median c. Standard deviation	b.	I by the extreme values? Mean None of the above	
3.	is the best measure of dispersion a. Mean c. Mean deviation	b.	Standard deviation None of the above	
4.	The best relative measure of dispersion isa. Coefficient of range		 Coefficient of Mean Deviation	

- 5. In a Poisson distribution with mean 4, then the variance is:
 - b. 4

a. 2 c. 0

d. None of the above

d. None of the above

- The number of girls in a family of four children, is an example of:
 - a. Binomial distribution

c. Coefficient of Variation

- b. Poisson distribution d. None of the above
- c. Normal distribution

a. t test c. F test

- b. Chi-square test d. Z test
- 8. If the calculated value of the test statistic is greater than its critical value, then:
 - a. The null hypothesis is not rejected

7. Degree of freedom is not associated with:

- b. The null hypothesis is rejected
- c. No conclusion
- d. None of the above
- 9. The two variables X and Y are independent, the correlation coefficient between X and Y is:
 - a. +1 c. ±1

b. -1

d. 0

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- 10. The product of two regression coefficients, is:
 - a. Less than or equal to 1
- b. Greater than or equal to 1

c. Equal to 1

d. None of the above

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llowing transposons is found in yeast?	
	a. Ty elementsc. Retroviral elements
s of almost 6 base pairs are called: b. Minisatellites	a. Microsatellite
d. Both b and c	c. VNTR
e is used by Relational database?	
b. JAVA	a. SQL
	c. Both a and b
tity in RDBMS is called an attribute?	Which of the entity in RI
b. Column	a. Row
d. All	c. Table
lowing file transfer protocol is used over remote network?	Which of the following f
b. SFTP	a. FTP
d. SSH	c. TCP
of haemophilusinfluenzae genome?	
b. 1800 bp	a. 1.8 kb c. Both
d. None	
r aligning translated nucleic acid 1 with translated nucleic acid 2?	Which tool is for alignin a. tblastx
b. tblastn d. All	c. Phi blast
llowing is a molecular structure database? b. MSD	which of the following i
d. None	c. Both
ptides will be least deflected by magnetic field in mass spectrometry	
b. 350 KDA	a. 120 KDA
d. 100 KDA	c. 1256 KDA
a feature of:	Global query is a feature
b. SRS	a. Entrez
d. None	c. Both

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Time: 2 hr. 30 mins.				
[Answer question no.1 & any four (4) from the rest]				
1.	a) Highlight the difference between eukaryotic and prokaryote genome with diagram.b) Write a note on secondary protein database.	5		
2.	a) How BLAST is performed? What is e value? b) What are the different types of BLAST?	5 6 4		
3.	 a) Define Population, Sample, Parameter and Statistic. b) A pharmaceutical company maintains that the mean time for a drug to take effect is 24 minutes. In a sample of 100 trials, the mean time is found to be 26 minutes with a standard deviation of 4 minutes. Can you say that the claim of the company is justified at 5% level of significance? [Given, the critical value of the test statistic at 5% significance level is 1.645] 	4		
4.	a) What is sequence homology? Write the steps of MSA? Which is the best method of phylogenetic alignment? Explain.b) Differentiate between motif and domain.			
5.	 a) Write the properties of Poisson distribution. b) If the probability of infection of COVD 19 is 20% in a certain place, what is the probability that out of 10 randomly selected people in the same place: i) Two are infected ii) At least three are infected 	4		
6.	Find mean, median, mode, standard deviation and coefficient of variation for the following distribution: Class: 10 - 20 20 - 30 30 - 40 40 - 50 50 - 60 Frequency: 5 8 12 16 18	10		
7.	Explain 2D gel electrophoresis with diagram. Write the significance of SDS in SDS PAGE.	10		
8.	 a) Differentiate between local and global alignment. b) Create PAM alignment for the following protein sequence AALVY ALLID 	5 5		

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